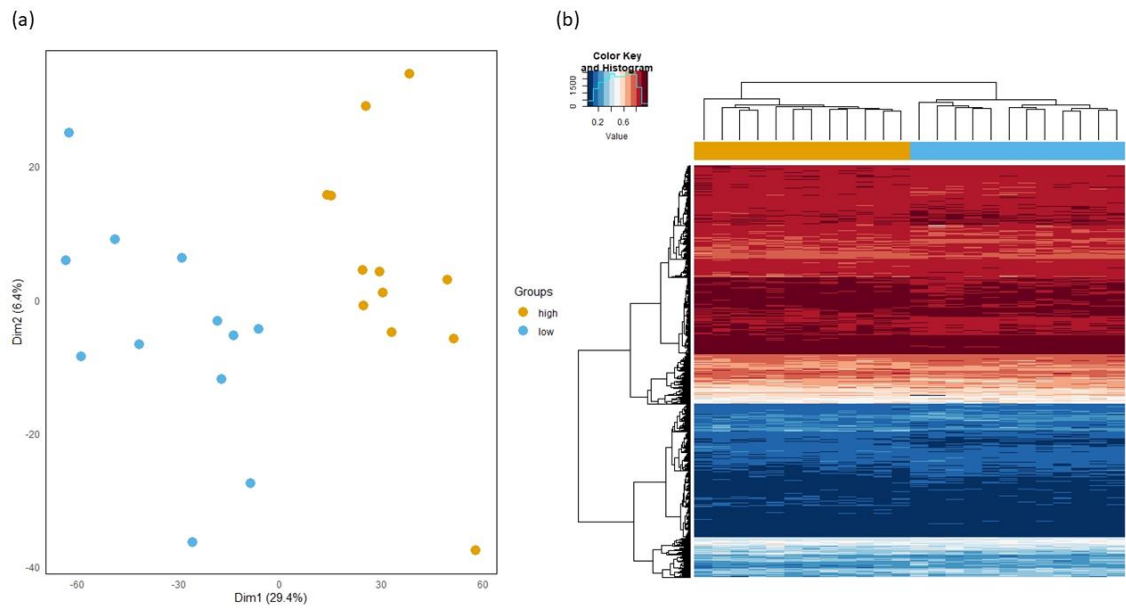


Figure 6



(a) Principal component analysis and (b) unsupervised clustering and heatmap of 5233 differentially methylated CpG sites detected using a threshold of an uncorrected p -value < 0.01 at 24 months of age and grouping by high and low vaccine responders.